

- 1 -  
piece 1, NC\_000913, yhiI\_yhiJ-, config: linear, direction: -, begin: 3629020, end: 3628606



- 2 -

```
{ p35-(23)-p10 3628801 Gap 1.4 bit  
{ p35-p10 3628801 total 6.3 bits
```

**p35** 5.3 bits ... p10

... p10

```
p35-(21)-p10 3628699 Gap 3.3 bits  
p35-p10 3628699 total 4.8 bits
```

**p10** 2.8 bits

[###> orf 38 codons

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```
{-----} sd-(18)-ir 3628625 Gap 6.9 bits  
|-----| sd-ir 3628625 yhiI_yhiJ- total 8.9 bits
```

p10 3.0 bits

```
{-----} p35-(25)-p10 3628628 Gap 4.0 bits  
|-----| p35-p10 3628628 total 4.3 bits
```

The diagram shows a sequence alignment between a nucleic acid-like sequence and a peptide sequence. The top sequence is **gt a a g c g c c a**, with the first letter 'g' underlined. The bottom sequence is **- val - ser - ala -**. Below the sequences are their corresponding labels: **\* 3628610** at the top right, and **5' - 3'** at the bottom left.

- lys - arg -

...  ir yhiI\_yhiJ-